RESINK ET AL.

Appl. No. 10/578,672

Atny Ref.: 3665-178

Amendment July 27, 2009

AMENDMENTS TO THE CLAIMS:

Please amend the claims as follows:

Claims 1-12 (Cancelled)

developing a Bovine Spongiform Encephalopathy (BSE) in a bovine or ovinean encaphalopathy in a mammal, comprising determining the presence, in a biological

13. (Currently Amended) A method for detecting the presence or the risk of

sample from the bovine or ovinemammal, of a target molecule selected in the group

consisting of:

a) a nucleic acid comprising a sequence of SEQ ID NO:1selected from SEQ ID

NO: 1-26 or a fragment thereof containing at least 5, preferably 6, 7, 8, 9 or 10

consecutive bases, and

b) a nucleic acid having a sequence complementary to a sequence according to

a),

c) a functional analogue of a nucleic acid according to a) or b) originating from

another species or a natural variant, or

d) a polypeptide coded by a nucleic acid according to a) to c),

the presence of said target molecule in the sample being an indication of the

presence or the risk of developing BSE an encephalopathy in said bovine or

ovinemammal.

14. (Previously Presented) Method according to claim 13, comprising

determining the presence of at least 2, 3, 4, 5, 6, 7, 8, 9, 10 or more target molecules.

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15. (Currently Amended) Method according to claim 13, comprising detecting

the presence or the absence of a nucleic acid according to a) to b)a) to c) by selective

hybridization or selective amplification.

Claim 16. (Cancelled)

Claim 17. (Cancelled)

18. (Currently Amended) Method according to claim 13, for detecting the

presence or the risk of developing BSE in a bovine or ovine, comprising contacting a

biological sample from the bovine or ovine containing nucleic acids with a product

comprising a support on which is immobilized at least one nucleic acid comprising the

sequence of SEQ ID NO:1a sequence selected from SEQ ID NO: 1-26, a fragment

thereof containing at least 5 consecutive bases, or a nucleic acid having a sequence

complementary thereto, and determining the hybridization profile, the profile indicating

the presence or the risk of developing BSE in the bovine or ovine.

19. (Previously Presented) Method according to claim 13, for detecting the

presence or the risk of developing BSE in a bovine or ovine, comprising contacting a

biological sample from the bovine or ovine containing nucleic acids with a product

comprising a primer allowing amplification of all or part of a target nucleic acid such as

defined in claim 13, said primer being single-stranded and having a length comprised

between 5 and 50 bases, the presence of amplification products being an indication of

the presence or the risk of developing BSE in the bovine or ovine.

20. (Previously Presented) The method of claim 13, wherein the sample is a

blood sample.

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Claims 21-23. (Cancelled)